

What is Multiple Alignment?

The MAO project is a joint effort to define community standards for data retrieval and exchange in the fields of DNA/RNA alignment, protein sequence and protein structure alignment.

Discuss Multiple Alignment Terminology

Please go to the [Multiple Alignment Discussion](#) page to leave comments regarding the Multiple Alignment Terminology.

Examples of comments may include:

- How you are using this terminology
- Why you decided to use (or not use) this terminology
- Strengths or limitations of the terminology
- Comparisons to other similar terminologies

Links

[Multiple Alignment FAQ](#)

[Multiple Alignment Home Page](#)